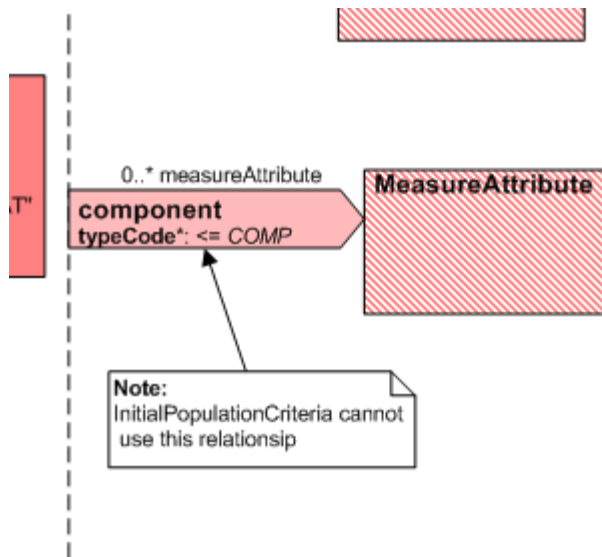


1. Please reopen this issue. Otherwise, we'll need to post a new one. This issue was posted on behalf of a team of people who are developing an application to consume HQMF eMeasures. It would help for the Assignee to communicate with us so a fully considered solution is provided.
2. The solution offered might work but it's not called out in the IG or the samples. Also, there's a note that the "COMP" relationship is not permitted for Initial Populations on the diagram in the standard. Given those contradictions, the solution offered in JIRA this doesn't appear to answer the need.



If it turns out that is the solution being offered, please explain the conflicting diagram note and omission from the QDM IG. An additional explanation would be helpful on why this different approach is used rather than the straightforward *title* and *text* elements.

3. An additional need is a way to group population sets together. All population sets appear at the same level and there is no *Organizer* segment. Is the serial position of populations (IPP, DENOM, NUMER ...) alone the only way to determine the association between the sets and groups?
4. In providing a solution, use the following eMeasure as an example:

EH_CMS171v4_NQF0527 SCIPInf1_AntibioticPriorIncision has 8 sets of populations. Use it as the "how to" example for organizing groups and distinguishing between the same population types.